

Does Not Comply
Corrected Diskette Needed



PCT09

RAW SEQUENCE LISTING

DATE: 10/01/2002

PATENT APPLICATION: US/09/913,770

TIME: 10:56:44

Input Set : A:\2590SEQ.txt

Output Set: N:\CRF4\10012002\I913770.raw

2 <110> APPLICANT: Takeda Chemical Industries, Ltd.
W--> 3 <120> TITLE OF INVENTION: Novel G Protein Coupled Receptor Protein and Its DNA
W--> 4 <130> FILE REFERENCE: 2590W00P
C--> 5 <140> CURRENT APPLICATION NUMBER: US/09/913,770
C--> 5 <141> CURRENT FILING DATE: 2001-08-17
Oh 5 <150> PRIOR APPLICATION NUMBER: JP 11-041336
6 <151> PRIOR FILING DATE: 1999-02-19
7 <150> PRIOR APPLICATION NUMBER: JP 11-125768
8 <151> PRIOR FILING DATE: 1999-05-06
W--> 9 <160> NUMBER OF SEQ ID: 4

ERRORED SEQUENCES

90 <210> SEQ ID NO: 4
91 <211> LENGTH: 25
92 <212> TYPE: DNA
93 <213> ORGANISM: Artificial Sequence
W--> 94 <220> FEATURE:
95 <223> OTHER INFORMATION:
W--> 96 <400> SEQUENCE: 4
C--> 97 ctaaaagtgt gatttcagag tg²⁵ttt
E--> 98 (1/4)

see items // an Error Summary Sheet

delete

↓
all bases

Must be in lower-case letters,
when using
new Sequence
Rules format

VERIFICATION SUMMARY

DATE: 10/01/2002

PATENT APPLICATION: US/09/913,770

TIME: 10:56:45

Input Set : A:\2590SEQ.txt

Output Set: N:\CRF4\10012002\I913770.raw

L:3 M:283 W: Missing Blank Line separator, <120> field identifier
L:4 M:283 W: Missing Blank Line separator, <130> field identifier
L:5 M:270 C: Current Application Number differs, Replaced Current Application No
L:5 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:9 M:283 W: Missing Blank Line separator, <160> field identifier
L:10 M:283 W: Missing Blank Line separator, <210> field identifier
L:14 M:283 W: Missing Blank Line separator, <400> field identifier
L:63 M:283 W: Missing Blank Line separator, <400> field identifier
L:64 M:112 C: (48) String data converted to lower case,
M:112 Repeated in SeqNo=2
L:86 M:283 W: Missing Blank Line separator, <220> field identifier
L:88 M:283 W: Missing Blank Line separator, <400> field identifier
L:89 M:112 C: (48) String data converted to lower case,
L:94 M:283 W: Missing Blank Line separator, <220> field identifier
L:96 M:283 W: Missing Blank Line separator, <400> field identifier
L:97 M:112 C: (48) String data converted to lower case,
L:98 M:254 E: No. of Bases conflict, LENGTH:Input:4 Counted:26 SEQ:4
L:98 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:98 M:252 E: No. of Seq. differs, <211> LENGTH:Input:25 Found:26 SEQ:4